

## POLYMORPHISMS IN THE FCER1A GENE

AAACAGAAGA	ATTAGTAAAG	GAATCCTGGA	GAAAGCCCCT	GCTGTGTATT	100
TAAAGGAGAA	AGGGAGATCA	TGTTGGGAAA	TTATAATATT	AAAAGTAAAC	
AAAAGCTAGG	AAGTAAATA	AAATAAATTA	TATGGCCTAG	ATCCCCATAA	200
GTAATGGTTT	AACTTCTGCC	TTCCTGTGTT	CTGAGCCAGA	TTAGGGCACA	
GTAGAGAAAG	AGGAGTCTCT	GAAAATGTTT	CCAATTTTCGC	TGGTCAGACA	300
GCGGATCATC	AGTGAATCAG	ATGAAAATTT	GTGGATTTAT	GCACTAACTG	
ATCAGCAGGA	AATTAACAA	GAAAAGCGTT	GGTAGCTCTG	GTGAATCCCA	400
AAAGAATTTG	GCAGTTGCTA	GCCATGCTCC	TGAATATGTA	TAAACAGTAC	
ATCATATGAC	TAAGAGTTTG	ACTTAGGGGT	TAGATTTTAT	GTGTTTGAAC	500
CCCAAATTAG	TTATTTAATA	GTTGGCACCC	CAAAACAAGT	TACTTAACCT	
CACTAAGATT	CAGTTTTTCT	GTTTATAAAA	TGTAGATAGT	GATAGTATGT	600
ACTTTATAGG	ATTATTGTGA	AAAATAAATG	AAATATCAGA	TTTATTTAGG	
G					
ATAACACCTG	GCATATGTTT	GGTATTCAGT	AATTAGTTGC	TGCTGTTTTA	700
TTCTGCTCTC	CCTTGCATCC	CACTTTTCTA	AGTTGTAAAC	TAAATAGTTG	
C					
TACACAGATT	GACAGATTAA	GAAAGGCTTG	TGATTGTGCT	AGACCTATGC	800
CTCTCTCTCA	CCAGATTCCA	GGTGTATATG	TGGAGGTGGG	ATAGGGAGTG	
GAGTAAGTGG	GTAAATATTA	AATTGCCAG	TTGGGCACCA	TCCTGAATAT	900
TATCTCTAAA	GAAAGAAGCA	AAACCAGGCA	CAGCTGATGG	GTTAACCAGA	
TATGATACAG	AAAACATTC	CTTCTGCTTT	TTGGTTTTAA	GCCTATATTT	
C T					
GAAGCCTTAG	ATCTCTCCAG	CACAGTAAGC	ACCAGGAGTC	CATGAAGAAG	1000
ATGGCTCCTG	CCATGGAATC	CCCTACTCTA	CTGTGTGTAG	CCTTACTGTT	
[exon 2: 1001..					
CTTCGGTAAG	TAGAGATTCA	ATTACCCCTC	CCAGGGAGGC	CCAAATGAAT	1100
A					
..1055]					
TTGGGGAGCA	GCTGGGGTAG	GAACCTTTAC	TGTGGGTGGT	GACTTTTTCT	1200
AGGACATGTG	CAAACATTG	GGCATTTCCT	AGGGACTCTG	TAGTGGAGCC	
AAGCTAGAAA	GCAGAGGCAA	GTGGGCTGAG	CAACACCTAA	GGAGGAAGCC	1300
AGACTGAAAG	CTTGGTTCCT	TGCATTTGCT	CTGGCATCTT	CCAGAGTGCA	
AATTTCTTAC	CAAGGTAATG	AGGGTAGAGG	AGAGAAAGAA	GCTCTTTCTT	1400
CCCCTGATTC	TCATTCTCTG	AAAGACGGTT	GGTCCTTAAA	ATTCCATGGA	
TGTAGATCTT	ATCCCCACAC	CCAGATTCTA	GTCCTCTGGA	GATAAAGAAG	1500
ACTGCTGGAC	ACTAATGTAT	CCTCTCTGGA	CTTTTGCAGC	TCCAGATGGC	
C A					
[exon 3: 1490..					
GTGTTAGCAG	GTGAGTCCTC	TGTTCTTGTT	CCCTTGGTGT	ATCAACATGT	
..1510]					
CTGGGCATTG	CTTTCCTCTC	ACTATTTTCT	TCGTCCCATC	ACTTCTGCTT	1600
TCTAATGAGC	ATGAATCTGT	TCCTTGGCCA	GACTACTTTC	CCTCTCCACC	
T					
TTGCCTTGTC	TTTCTTTTTT	TCCCTGATTC	ATTGCATTCT	CTCAAGTCAT	1700
TCTCTCCTCT	GTTTTAGTCA	ATAACCATGT	CTGTTGCACA	TATACATGTC	
TCATTCTCTC	TCCTAGACAC	TTTGGCATGA	TCTCGCTCAA	TAATTACATT	1800
ATTATTATTA	TTGCCATTTT	ATAATTGAGG	ATGCTGAAAC	TCAGTGATTT	
TCTGGTGGTT	ACATGGCTAA	GGAAGTGGAT	TTCAACGTAA	GTTCTTGGGA	1900
TCTAAGTCCA	GTTCTCTTCT	GACTATATCA	CCCTTTTGTT	ATCACCATGT	
ATCTACTTCT	TTGGTCTCTG	TTCAAATTTG	CACTACATCC	CCTTGTTCCA	2000
GGAAGCCATT	CAAGACTGAC	TTTCTTAGTG	CCTCTCACTA	CTTCTGGAA	

FIGURE 1A

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CTGACATATG	TTTTTCACTC	TGTATATACT	TACAATTAAA	TAGTCATAAA	2100
TATTCAGAGC	TTGGAGAAAC	CTTATATTTT	ATCCAGTCCA	GTAAATTTAT	
CCATCCATAA	TTCACCTCATT	CATTACACATA	ATAAATATTT	AATGTAACAA	2200
TGGTTGAACA	TGGCAGACAG	TGTTTCTACC	TCAAAAGAGA	TTGCAGTCCT	
CATTTACAGA	TACTGAATTG	AAATTAACAG	AAGTAGAGTG	AGTCAGCTCA	2300
AATCACATAG	TGAATTGGTT	TCTTTGTTTT	TAAATCTCCT	GCATATGTGT	
CCTGTCTTTC	TCCCTGTGTT	GGGCGTTCCC	TGGGGCACCA	ATACTAATTT	2400
CTCCTTCCCC	TAGAAATCAA	AACAGGGTCT	TATCACCAAC	AGAATAAGGA	
	G				
CAGGTTGACC	ACTGATTGTC	AGAATATTGC	TTCGTTTGTA	CTTTTAAGCC	2500
TAGACAGTTT	TCAATGACTT	TTTTTCTCTC	TACATGTCTT	TTCATATTTT	
TATCTTCTTG	AAGTCCCTCA	GAAACCTAAG	GTCTCCTTGA	ACCCTCCATG	2600
[exon 4: 2564..					
GAATAGAATA	TTTAAAGGAG	AGAATGTGAC	TCTTACATGT	AATGGGAACA	
ATTTCTTTGA	AGTCAGTTCC	ACCAAATGGT	TCCACAATGG	CAGCCTTTCA	2700
GAAGAGACAA	ATTCAAGTTT	GAATATTGTG	AATGCCAAAT	TTGAAGACAG	
		G			
TGGAGAATAC	AAATGTCAGC	ACCAACAAGT	TAATGAGAGT	GAACCTGTGT	2800
			A		
ACCTGGAAGT	CTTCAGTGGT	AAGTTCCAGG	GATATGGAAA	TACAGATCTC	
	..2818]				
TCATGTGAGG	GATGGCTCAT	CTGAAGATGG	GAAAAACAG	GTTATTCCAA	2900
GGGTTAGGAC	ACCAGAGTGG	GATTCAAGGC	CTCTCATTTT	TAAGACCCCT	
		C			
GCATTGGCTG	GGCACAGTGG	CTCACGCCTG	TAATCCCAGC	ACTTTGGGAG	3000
			A		
GCTGAGGCAG	GTGGATCAGC	AGGTCAGGAG	ATCGAGACCA	TCCGGCTAAC	
			A		
ATGGTGAAAC	CCCATCTCTG	CTAAAAAATA	TATATATATA	AAATTAGCCG	3100
GGCGTAGTGG	TGGGCACCTG	TAGTCCCAGG	TACTCGGGAG	GCTGAGGCAG	
GAGAATGGTG	TGAACCCAGG	AGGTGGAGGT	TGCAGTGAGC	TGAGATCACG	3200
CCACTGCCCT	CCAGCCTGGG	CTACAGAGCA	AGACTCCGTC	TCAAAAAATA	
AATAAATAAA	TAAAAAAGAC	CCCTGCATCT	CTTTTCTTCT	ACCCCTTCC	3300
CTTTTGATTA	CTTGATGACC	TTCTTTCAAT	ATTCTAGTCA	TCTCTCAATA	
TTATTCCTCC	ACCCTATTTT	CCTCTATCTT	TTCTGCCTAG	ATTCAGGTAT	3400
ATATTATGTG	GTCAAACAGC	ATGACATATA	TGTGAACATT	TCAAAGAGCT	
GTGTATCTGG	AATAGGATCA	AAAGGTTTGA	CTTAAAGTTT	TGCTCTGCAT	3500
AATCCATATG	GCAGGACCTG	AATATTAGGT	TGTACTCTTC	GTTATGAAAC	
ATATCTGGGT	ACATTTCTTT	ATGTCCTCTG	TTGTTACTTA	AGAACACATA	3600
TTTCATGCTT	GTTTCATTTT	TATCACTCCT	ACTGCCAACA	AATAGCATAG	
CATGCTTAGG	CACATGTGGC	TTAATTAGCA	AATGTTGAAT	AAACAAATTA	3700
ATGATTTTGA	ATAGTGACCA	ATAGGTCTCT	TTTATACTCT	ATATTTTCT	
CTTGAGTGAA	AAAAAATGTT	TCAACCTCCA	TATGTAAATT	CCAAACACAA	3800
ACTAAAGCAA	TGTAGAATAG	CTTCTTTATT	CCCTGGAGTA	GGTTCTAGAG	
AAGTCCTAAA	GGATTGGTCC	TAAATTAATT	ATGCTTATTA	TGCTAGCGAT	3900
ATTTCCCTTT	AAAATTCTCC	TTTAATGAAT	GCTTTTAAAT	TTTTACAAAA	
GCATTAACCA	TAGAATGTGA	TTCTTGCTCT	TCACTGACTC	ATTAGTGACA	4000
AATATTTGTT	GAGTACCTAC	CAACTCCTAA	GTATTGCTAC	CAACTCCTAA	
ATACTGTGTT	GGGCATTGAG	AATAGAATGT	AGAACTAGAC	AGGGTCCCTG	4100
ACTTCTTGGA	GCACAGAGCA	GTATGGGAAG	AGGACATTAA	ATAAAGAATT	
ACATAAGTAA	TTAATTTAAA	TTATACATGT	TTTGAAGAAG	TTTTTTTTTG	4200
ACAACTATAA	TTAACACTAG	AACTGGGAAG	TTTCTATAAG	GTAAGAGAGG	
ACAAAATAGA	CACTCTCCTA	AGCTAAAATT	CCCAAGAAAG	ACTGTTTATT	4300
TTCCCCTAAC	TAAC TAGAAC	TAGCAACAGA	AGATCTGAAA	GGAATTCTGG	

FIGURE 1B

CTTTCAAGTG TTCCATGTAT GGACTCATCA GGGAGGTCCG AGAGGCTTTG 4400  
TGGCCCCAGA CTGACTTTTC AGGAGGGGAA AGGATTTATC AATACACAAG  
ACAGGCTCTA AGCATTATTT TGTGCCCTTT AAAAATCCAC TTTATGAGCC 4500  
AAAAAGTGAG TTAATGATAA TTCATAGTTT CTGACACATG CTCTATGCGT  
GGCTCTCTTT TCTCTATTCA TTCTCTCTCT CTTCATTTAT TGTTAAATAA 4600  
A  
ATAATGTAAT GAATGTTCTT CAGACTGGCT GCTCCTTCAG GCCTCTGCTG  
[exon 5: 4624..  
AGGTGGTGAT GGAGGGCCAG CCCCTCTTCC TCAGGTGCCA TGGTTGGAGG 4700  
AACTGGGATG TGTACAAGGT GATCTATTAT AAGGATGGTG AAGCTCTCAA  
GTACTGGTAT GAGAACCACA ACATCTCCAT TACAAATGCC ACAGTTGAAG 4800  
ACAGTGGAAC CTACTACTGT ACGGGCAAAG TGTGGCAGCT GGACTATGAG  
T  
TCTGAGCCCC TCAACATTAC TGTAATAAAA GGTGAGTTGG TAAAGGAAAG 4900  
..4881]  
GAAAAGCATC CATAGCAGGG GAAGGAAGAG AGAACTTCTG AGCCTGAGCA  
GTTGCAGCTT GTAGAAGGGG GGCACCTGTG ATACACTGGA AAGCCTACCA 5000  
T  
GACTTGCAAT GAGGAGACCT GGGTGATAGT ATATATCTCA ATCTCTGTTT  
CAAAGCCTTG ACTTGTTAAA TGGTGATAGT AATACCTGCT TGCATATGA 5100  
C  
AATTTTTATG AAGATTAATG TGGTAATATT TGTGAAATGA CTTTGTAAC  
TGTTAAGCAC TACCCAAGCA TAACAGATTG TGATTACTAT TTTGATCTCA 5200  
AAGTCATCTG TTGCTCCTGG GGAACACTT ATATTTATCA AATTGAAAAA  
AAGTTTCAAA GTTGAATGAA GAAAGGATAT AAAGAGCTTG AGGAGCCCAT 5300  
TCCAGCTTAG GAGGGCTGGG AAAGGAAACC AGCAAGTCAG TAAGCTGTGT  
GCCTGTGTAT TGAGGGAGGA GGAATGGAC TTGATATGGA GAGGGTAGGG 5400  
AGGTGGACTG CCTCTATGGC CTGTAAGAAA AACTGCTCTC TCCAAACTCT  
TTATAAGAGA GGGAGCCTGT GAAGTATTCA CTTTTGAAGG AGAAAGTTAG 5500  
ACTTTTCCTT CACACACTTT GTACATAATA ATGTTTAAAA AAGCATGAGG  
TCAAAATACA TAATTAAGTC CTAGCAGTTC TCTGTTAACT AATTTGAGAC 5600  
TGAAGTGCTA TGTACTTGTC TCTAGGCTTC CAGTATCTTC ATCTGTAAAA  
CAGAATATTT GGTCTAGATT CCATTAGAAT CATTTGATAA CTTAAAAAAT 5700  
ATATTGATGC TCATGTCTCA TTTCTTGAGA TTCTGATTTA ATGGGTTTGG  
GGTGCAGCCT GGGTATACGT ATTTTTCATA GGTCTTTCAC ATAATGGTAA 5800  
TGGGTAGCCA ATATTGAGAA TCACTTGTCT AGGTGATCTT TAAATGATTT  
CTGGATGTAA TATTCTGAGG CTCTATAATT TGAGACTAAT CACAAAAATC 5900  
GGTACAGTTT ATAAACAGAC TAACAGAACC ACAAATAAT AGAATTGGAA  
GGCAATTTAA CTAGTGCAAT TTCTTCATTT TGCCTAACAG GCATGTAAGA 6000  
AATGATGATT GATTGAGTAA TAGGCATTGA TGACCCCTGT CCTCACTTTG  
TCCCCTTTCC ACCCCTTAAT TATATGTGAA TTCTGGTCTT GTCATTTCGA 6100  
ATAAGGGGTT TATCTTTCCT ATTGTCTTCC CCTCTGGGCA CGGCACACTG  
GCTACTGGAG TTAAGAGGAA ATGCTTAGGA CTCCCTGTGG CTCCAGGGAG 6200  
CACCAACAGA GCAACTCAAC CTAGTGTTAA TCTGAGTGTT TTCTCTGTGC  
TTCTGGATGC CACATCACGC TAAAAATGAA GGACAAAGCT TGGTCTTTCT 6300  
CTTAGGGAGG ATGAAACTCT GAACCTCATT TTTCAGTTCC CAAGATGAAT  
TATGTTTCTC ATTGCATCTG TGTTCCACTA CAGCTCCGCG TGAGAAGTAC 6400  
[exon 6: 6384..  
TGGCTACAAT TTTTATATCC ATTGTTGGTG GTGATTCTGT TTGCTGTGGA  
CACAGGATTA TTTATCTCAA CTCAGCAGCA GGTACATTT CTCTTGAAGA 6500  
TTAAGAGAAC CAGGAAAGGC TTCAGACTTC TGAACCCACA TCCTAAGCCA  
A  
AACCCCAAAA ACAACTGATA TAATTACTCA AGAAATATTT GCAACATTAG 6600  
..6568]

FIGURE 1C

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TTTTTTTCCA	GCATCAGCAA	TTGCTACTCA	ATTGTCAAAC	ACAGCTTGCA	
		C		G	
ATATACATAG	AAACGTCTGT	GCTCAAGGAT	TTATAGAAAT	GCTTCATTAA	6700
ACTGAGTGAA	ACTGGTTAAG	TGGCATGTAA	TAGTAAGTGC	TCAATTAACA	
	A				
TTGGTTGAAT	AAATGAGAGA	ATGAATAGAT	TCATTTATTA	GCATTTGTAA	6800
AAGAGATGTT	CAATTTCAAT	AAAATAAATA	TAAAACCATG	TAACAGAATG	
CTTCTGAGTA	TTCAAGGCTT	GCTAGTTTGT	TTGTTTGTTT	TCTACTAAAG	6900
GCAAGGACCA	TGAAGTTCTA	GATTGGAAAT	GTCCTCTCTT	GAATATTGCA	
AGTGCGATCT	AGGAATGAAA	AGACATAGGA	GGATGCCAGT	GAGGTGGATC	7000
ATTTTATGTC	TTCTTCTTCA	GCTTACTAAA	TATGAACTTT	CAGTTCTTGG	
CAGAATCAGG	GACAGTCTCA	AGACATAGGA	CTCTCAGGAT	GAAGTAGAGT	7100
CCAGGATTCC	TCTGTGATTG	TTTTGCCCCCT	CCCAAATTTA	TATCTTGAAC	
TTATGTCTTG	TATCTTTATA	CAGCACCTGA	ACCAAGCATT	TTGGAGAAAT	7200
TCCAGCTAAT	AATAATAACC	AAAACCTTCG	GCTCTGAAAA	CAGTCCAGGA	
CTGAATAAGA	TCTTGGGCAA	AAGAACTAGA	CAGTTTTGGT	TTATTTTCCC	7300
TTTCATTTTA	TGTCTTCATC	ATAGTCATTG	GAGGCTCATT	CTTCTTGTC	
TGGAGTAAAT	GGGATTAAAG	TT			7372

FIGURE 1D

## POLYMORPHISMS IN THE CODING SEQUENCE OF FCER1A

ATGGCTCCTG	CCATGGAATC	CCCTACTCTA	CTGTGTGTAG	CCTTACTGTT	
CTTCGCTCCA	GATGGCGTGT	TAGCAGTCCC	TCAGAAACCT	AAGGTCTCCT	100
TGAACCCTCC	ATGGAATAGA	ATATTTAAAG	GAGAGAATGT	GA CTCTTACA	
TGTAATGGGA	ACAATTTCTT	TGAAGTCAGT	TCCACCAAAT	GGTTCCACAA	200
TGGCAGCCTT	TCAGAAGAGA	CAAATTCAAG	TTTGAATATT	GTGAATGCCA	
AATTTGAAGA	CAGTGGAGAA	TACAAATGTC	AGCACCAACA	AGTTAATGAG	300
G					
AGTGAACCTG	TGTACCTGGA	AGTCTTCAGT	GA CTGGCTGC	TCCTTCAGGC	
A					
CTCTGCTGAG	GTGGTGATGG	AGGGCCAGCC	CCTCTTCCTC	AGGTGCCATG	400
GTTGGAGGAA	CTGGGATGTG	TACAAGGTGA	TCTATTATAA	GGATGGTGAA	
GCTCTCAAGT	ACTGGTATGA	GAACCACAAC	ATCTCCATTA	CAAATGCCAC	500
AGTTGAAGAC	AGTGAACCT	ACTACTGTAC	GGGCAAAGTG	TGGCAGCTGG	
		T			
ACTATGAGTC	TGAGCCCCTC	AACATTACTG	TAATAAAAGC	TCCGCGTGAG	600
AAGTACTGGC	TACAATTTTT	TATCCCATTG	TTGGTGGTGA	TTCTGTTTGC	
TGTGGACACA	GGATTATTTA	TCTCAACTCA	GCAGCAGGTC	ACATTCTCT	700
TGAAGATTAA	GAGAACCAGG	AAAGGCTTCA	GA CTTCTGAA	CCCACATCCT	
			A		
AAGCCAAACC	CCAAAAACAA	CTGA			774

FIGURE 2

## ISOFORMS OF THE FCER1A PROTEIN

MAPAMESPTL	LCVALLFFAP	DGVLAVPQKP	KVSLNPPWNR	IFKGENVTLT	
CNGNNFFEVS	STKWFHNGSL	SEETNSSLNI	VNAKFEDSGE	YKCQHQQVNE	100
			R		
SEPVYLEVFS	DWLLLQASAE	VVMEGQPLFL	RCHGWRNWDV	YKVIYYKDGE	
N					
ALKYWYENHN	ISITNATVED	SGTYICTGKV	WQLDYESEPL	NITVIKAPRE	200
		M			
KYWLQFFIPL	LVVILFAVDT	GLFISTQQQV	TFLKIKRTR	KGFRLLNPHP	
				K	
KPNPKNN					257

ISOFORMS OF THE FCER1A GENE

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Contact: Gisela Field  
Chew et al.  
203-786-3473

FIGURE 3